```
SEQUENCE LISTING
```

10/500184

```
<110> SUGO, Izumi
TOMONOU, Kikuo
```

<120> METHOD FOR STABILIZING PROTEINS

<130> 14875-132US1

<140> US 10/500,184

<141> 2004-06-25

<150> PCT/JP02/13804

<151> 2002-12-27

<150> JP 2001-400895

<151> 2001-12-28

<160> 28

<170> PatentIn Ver. 2.1

<210> 1

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:an artificially synthesized primer sequence

<400> 1

aattggaagc ttgc

14

<210> 2

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<400> 2

ccttcgaacg ttaa

14

<210> 3

<211> 41

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:an artificially synthesized primer sequence

<400> 3 gagtctagaa tggattggtg ggaatgatcc tgcgaatatg c	41
<210> 4 <211> 40 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 4 gagaatttcg ggtcatacat actatgcata ttcgcaggat	40
<210> 5 <211> 43 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 5 gagtctagaa tggattggtg ggaatgatcc tgcgaataag cat	43
<210> 6 <211> 40 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 6 gagaatttcg ggtcatacat actatgctta ttcgcaggat	40
<210> 7 <211> 43 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 7 gagtctagaa tggattggtg ggaatgatcc tgcgaattgg cat	43

<210> 8

```
<211> 40
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
gagaatttcg ggtcatacat actatgccaa ttcgcaggat
                                                                    40
<210> 9
<211> 43
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
gagtctagaa tggattggtg ggaatgatcc tgcgaatcag cat
                                                                    43
<210> 10
<211> 40
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
gagaatttcg ggtcatacat actatgctga ttcgcaggat
                                                                    40
<210> 11
<211> 43
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: an artificially
      synthesized primer sequence
gagtctagaa tggattggtg ggaatgatcc tgcgaatgag cat
                                                                    43
<210> 12
<211> 40
<212> DNA
<213> Artificial Sequence
<220>
```

<223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 12 gagaatttcg ggtcatacat actatgctca ttcgcaggat	40
<210> 13 <211> 43 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 13 gagtctagaa tggattggtg ggaatgatcc tgcgaatttc cat	43
<210> 14 <211> 40 <212> DNA <213> Artificial Sequence	i
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 14 gagaatttcg ggtcatacat actatggaaa ttcgcaggat	40
<210> 15 <211> 43 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 15 gagtctagaa tggattggtg ggaatgatcc tgcgaatacc cat	43
<210> 16 <211> 40 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> 16 gagaatttcg ggtcatacat actatgggta ttcgcaggat	4.0

<210><211><211><212><213>	43	
<220> <223>	Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> gagtc	17 tagaa tggattggtg ggaatgatcc tgcgaataac cat	43
<210><211><211><212><213>	40	
<220> <223>	Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> gagaat	18 tttcg ggtcatacat actatggtta ttcgcaggat	40
<210><211><211><212><213>	43	
<220> <223>	Description of Artificial Sequence:an artificially synthesized primer sequence	
<400> gagtct	19 cagaa tggattggtg ggaatgatcc tgcgaatgac cat	43
<210><211><211><212><213>	40	
<220> <223>	Description of Artificial Sequence: an artificially synthesized primer sequence	
<400> gagaat	20 Ettcg ggtcatacat actatggtca ttcgcaggat	40
<210>		

<212> DNA

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
                                                               . 43
gagtctagaa tggattggtg ggaatgatcc tgcgaatccc cat
<210> 22
<211> 40
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
gagaatttcg ggtcatacat actatgggga ttcgcaggat
                                                                    40
<210> 23
<211> 43
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
<400> 23
gagtctagaa tggattggtg ggaatgatcc tgcgaattgc cat
                                                                    43
<210> 24
<211> 40
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:an artificially
      synthesized primer sequence
gagaatttcg ggtcatacat actatggcaa ttcgcaggat
                                                                    40
<210> 25
<211> 444
<212> PRT
<213> Homo sapiens
<400> 25
Gln Val Gln Leu Leu Glu Ser Gly Ala Val Leu Ala Arg Pro Gly Thr
                                     10
```

- Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Phe Asn Ile Lys Asp Tyr 20 25 30
- Tyr Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile 35 40 45
- Gly Gly Asn Asp Pro Ala Asn Gly His Ser Met Tyr Asp Pro Lys Phe
 50 60
- Gln Gly Arg Val Thr Ile Thr Ala Asp Thr Ser Thr Ser Thr Val Phe
 65 70 75 80
- Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95
- Ala Arg Asp Ser Gly Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Leu 100 105 110
- Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu 115 120 125
- Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys 130 135 140
- Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser 145 150 155 160
- Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser 165 170 175
- Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser 180 185 190
- Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser Asn 195 200 205
- Thr Lys Val Asp Lys Arg Val Glu Ser Lys Tyr Gly Pro Pro Cys Pro 210 215 220
- Pro Cys Pro Ala Pro Glu Phe Leu Gly Gly Pro Ser Val Phe Leu Phe 225 230 235 240
- Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val 245 250 255
- Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Glu Val Gln Phe 260 265 270
- Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro 275 280 285
- Arg Glu Glu Gln Phe Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr 290 295 300
- Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val 305 310 315 320

Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu Lys Thr Ile Ser Lys Ala 325 330 335

Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Gln 340 345 350

Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly 355 360 365

Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro 370 380

Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser 385 390 395 400

Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp Lys Ser Arg Trp Gln Glu 405 410 415

Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His
420 425 430

Tyr Thr Gln Lys Ser Leu Ser Leu Ser Leu Gly Lys
435
440

<210> 26

<211> 214

<212> PRT

<213> Homo sapiens

<400> 26

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Asp Ile Lys Ser Phe 20 25 30

Leu Ser Trp Tyr Gln Gln Lys Pro Glu Lys Ala Pro Lys Ser Leu Ile 35 40 45

Tyr Tyr Ala Thr Ser Leu Ala Asp Gly Val Pro Ser Arg Phe Ser Gly 50 55 60

Ser Gly Ser Gly Thr Asp Tyr Thr Leu Thr Ile Ser Ser Leu Gln Pro 65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Leu Gln His Gly Glu Ser Pro Tyr 85 90 95

Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala
100 105 110

Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
115 120 125

Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala

Lys V 145	/al Gln	Trp	Lys	Val 150	Asp	Asn	Ala	Leu	Gln 155	Ser	Gly	Asn	Ser	Gln 160	
Glu S	Ser Val	Thr	Glu 165	Gln	Asp	Ser	Lys	Asp 170	Ser	Thr	Tyr	Ser	Leu 175	Ser	
Ser T	Thr Leu	Thr 180	Leu	Ser	Lys	Ala	Asp 185	Tyr	Glu	Lys	His	Lys 190	Val	Tyr	
Ala C	Cys Glu 195	Val	Thr	His	Gln	Gly 200	Leu	Ser	Ser	Pro	Val 205	Thr	Lys	Ser	
	Asn Arg	Gly	Glu	Cys											
<210><211><211><212><213>	37	icial	l Sec	quenc	ce					•					
<220> <223> Description of Artificial Sequence:an artificially synthesized sequence															
<400> 27 gagtctagaa tggattggtg ggaatgatcc tgcgaat												37			
<210><211><211><212><213>	39	icial	l Seq	quenc	ce										
<220> <223> Description of Artificial Sequence:an artificially synthesized sequence															
<222>	misc_f (1) n = g,	(2)		or t											
<400> 28 nnattcgcag gatcattccc accaatccat tctagactc												39			